

## Sequence PC19084 12192003.ST25.txt SEQUENCE LISTING

<110> Agouron Pharmaceuticals, Inc./A Pfizer Company

<120> PIN1 PEPTIDYL-PROLYL ISOMERASE POLYPEPTIDES, THEIR CRYSTAL STRUCTURES, AND USE THEREOF FOR DRUG DESIGN

PC19084A <130> 10/616,003 <140> 2003-07-08 <141> us 60/394,889 <150> <151> 2002-07-09 <160> 11 <170> PatentIn version 3.1 <210> 423 <211> <212> DNA Artificial <213> <220> **PPlase** <223> <400> atgggcagca gccatcatca tcatcatcac agcagcggcc tggtgccgcg cggcagccat 60 atgggcaaaa acgggcaggg ggagcctgcc agggtccgct gctcgcacct gctggtgaag 120 cacagccagt cacggcggcc ctcgtcctgg cggcaggaga agatcacccg gaccaaggag 180 gaggccctgg agctgatcaa cggctacatc cagaagatca agtcgggaga ggaggacttt 240 300 gagtctctgg cctcacagtt cagcgactgc agctcagcca aggccagggg agacctgggt 360 gccttcagca gaggtcagat gcagaagcca tttgaagacg cctcgtttgc gctgcggacg 420 ggggagatga gcgggcccgt gttcacggat tccggcatcc acatcatcct ccgcactgag 423 tga <210> 2 123 <211> <212> PRT Artificial <213> <220> <223> **PPlase** <400> 2

Gly Ser His Met Gly Lys Asn Gly Gln Gly Glu Pro Ala Arg Val Arg

Cys Ser His Leu Leu Val Lys His Ser Gln Ser Arg Arg Pro Ser Ser 20 25 30

Sequence PC19084 12192003.ST25.txt Trp Arg Gln Glu Lys Ile Thr Arg Thr Lys Glu Glu Ala Leu Glu Leu Ile Asn Gly Tyr Ile Gln Lys Ile Lys Ser Gly Glu Glu Asp Phe Glu 50 60 Ser Leu Ala Ser Gln Phe Ser Asp Cys Ser Ser Ala Lys Ala Arg Gly 65 70 75 80 Asp Leu Gly Ala Phe Ser Arg Gly Gln Met Gln Lys Pro Phe Glu Asp Ala Ser Phe Ala Leu Arg Thr Gly Glu Met Ser Gly Pro Val Phe Thr 100 Asp Ser Gly Ile His Ile Ile Leu Arg Thr Glu 115 120 <210> 422 <211> <212> DNA Artificial <213> <220> <223> **PPlase** <400> atgggcagca gccatcatca tcatcatcac agcagcggcc tggtgccgcg cggcagccat 60 atggcaaaaa cgggcagggg gagcctgcca gggtccgctg ctcgcacctg ctggtgaagc 120 180 acagccagtc acggcggccc tcgtcctggc ggcaggagca gatcacccgg acccaggagg aggccctgga gctgatcaac ggctacatcc agaagatcaa gtcgggagag gaggactttg 240 agtctctggc ctcacagttc agcgactgca gctcagccaa ggccagggga gacctgggtg 300 360 ccttcagcag aggtcagatg cagaagccat ttgaagacgc ctcgtttgcg ctgcggacgg gggagatgag cgggcccgtg ttcacggatt ccggcatcca catcatcctc cgcactgagt 420 422 ga <210> 123 <212> PRT Artificial <213> <220> <223> **PPlase** <400> Gly Ser His Met Gly Lys Asn Gly Gln Gly Glu Pro Ala Arg Val Arg
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## Sequence PC19084 12192003.ST25.txt Cys Ser His Leu Leu Val Lys His Ser Gln Ser Arg Arg Pro Ser Ser 20 25 30 20 Trp Arg Gln Glu Gln Ile Thr Arg Thr Gln Glu Glu Ala Leu Glu Leu 35 40 45 Ile Asn Gly Tyr Ile Gln Lys Ile Lys Ser Gly Glu Glu Asp Phe Glu 50 60Ser Leu Ala Ser Gln Phe Ser Asp Cys Ser Ser Ala Lys Ala Arg Gly 65 70 75 80 Asp Leu Gly Ala Phe Ser Arg Gly Gln Met Gln Lys Pro Phe Glu Asp Ala Ser Phe Ala Leu Arg Thr Gly Glu Met Ser Gly Pro Val Phe Thr Asp Ser Gly Ile His Ile Ile Leu Arg Thr Glu 115 120 <210> <211> 36 <212> DNA Artificial <213> <220> <223> Primer <400> 36 agcagccata tgggcaaaaa cgggcagggg gagcct <210> 30 <211> <212> DNA Artificial <213> <220> <223> Primer <400> 6 30 cttggatcct cactcagtgc ggaggatgat <210> 7 119 <211> <212> PRT Artificial <213>

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7

PPlase domain

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Sequence PC19084 12192003.ST25.txt
Gly Lys Asn Gly Gln Gly Glu Pro Ala Arg Val Arg Cys Ser His Leu
1 10 15
Leu Val Lys His Ser Gln Ser Arg Arg Pro Ser Ser Trp Arg Gln Glu 20 25 30
Lys Ile Thr Arg Thr Lys Glu Glu Ala Leu Glu Leu Ile Asn Gly Tyr 35 40
Ile Gln Lys Ile Lys Ser Gly Glu Glu Asp Phe Glu Ser Leu Ala Ser 50 60
Gln Phe Ser Asp Cys Ser Ser Ala Lys Ala Arg Gly Asp Leu Gly Ala 65 70 75 80
Phe Ser Arg Gly Gln Met Gln Lys Pro Phe Glu Asp Ala Ser Phe Ala
Leu Arg Thr Gly Glu Met Ser Gly Pro Val Phe Thr Asp Ser Gly Ile 100 105 110
His Ile Ile Leu Arg Thr Glu
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       44
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       44
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                                                                             44
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<223> PPlase domain
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## Sequence PC19084 12192003.ST25.txt

<400> 10

Gly Lys Asn Gly Gln Gly Glu Pro Ala Arg Val Arg Cys Ser His Leu  $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$ 

Leu Val Lys His Ser Gln Ser Arg Arg Pro Ser Ser Trp Arg Gln Glu 20 25 30

Gln Ile Thr Arg Thr Gln Glu Glu Ala Leu Glu Leu Ile Asn Gly Tyr 35 40 45

Ile Gln Lys Ile Lys Ser Gly Glu Glu Asp Phe Glu Ser Leu Ala Ser 50 55 60

Gln Phe Ser Asp Cys Ser Ser Ala Lys Ala Arg Gly Asp Leu Gly Ala 65 70 75 80

Phe Ser Arg Gly Gln Met Gln Lys Pro Phe Glu Asp Ala Ser Phe Ala 85 90 95

Leu Arg Thr Gly Glu Met Ser Gly Pro Val Phe Thr Asp Ser Gly Ile  $100 \hspace{1cm} 105 \hspace{1cm} 110$ 

His Ile Ile Leu Arg Thr Glu 115

<210> 11

<211> 11

<212> PRT <213> Artificial

<220>

<223> Pintide where the serine is a phosphorylated

<400> 11

Phe Leu Trp Phe Tyr Pro Ser Pro Phe Leu Glu 1 5 10